RAW SEQUENCE LISTING

The Biotechnology Systems Branch of the Scientific and Technical Information Center (STIC) no errors detected.

Application Serial Number:	<u> (0/55</u> 1, 699
Source:	PCT
Date Processed by STIC:	10/14/2005
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PCT

RAW SEQUENCE LISTING DATE: 10/14/2005
PATENT APPLICATION: US/10/551,699 TIME: 13:27:41

Input Set : A:\CD054PCT.ST25.txt

Output Set: N:\CRF4\10142005\J551699.raw

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3 <110> APPLICANT: CropDesign N.V.
      5 <120> TITLE OF INVENTION: Plant haemoglobin
      7 <130> FILE REFERENCE: CD-054-PCT
C--> 9 <140> CURRENT APPLICATION NUMBER: US/10/551,699
C--> 9 <141> CURRENT FILING DATE: 2005-09-30
     9 <150> PRIOR APPLICATION NUMBER: EP 03075974.0
    10 <151> PRIOR FILING DATE: 2003-04-01
    12 <160> NUMBER OF SEQ ID NOS: 12
    14 <170> SOFTWARE: PatentIn version 3.1
    16 <210> SEQ ID NO: 1
    17 <211> LENGTH: 860
    18 <212> TYPE: DNA
    19 <213> ORGANISM: Beta vulgaris
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    24 taqtttcttt tcttttaaat taaaacaaaa ctatqacttt tacaqaqaaa gatqaagctt
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    26 tggtaaaaga atcatgggat ataatgaagc aaaatatccc agaatacagc cttcggtttt
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    28 tetecataat attggaaatt getecageag ecaaaaatat gtteteattt ttaagggatt
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    30 cagaggaagt tccacagaat aatcccaagc tgaaagctca tgcaatcaag gtttttaaaa
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    32 tgacatgtga atcagccatt caacttcgag aaaaaggtga agtggttgta ggagagacta
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    34 cccttaaata tttgggagct atccatttga agaatggagt gattgatccc cattttgagg
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    36 ttgtgaaaca agcattattg agaaccatag aagaagcaag tggtgacaaa tggagtgaag
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    40 tgaaggaata ggtagcttag ttctcagtcg ccaaaagtat tactctaaaa atattgaata
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    42 aatattetta ttgtttttga ggggaaatta ttgttattgt tgattetgae teaettattt
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    44 atccgagtga cttgatatgg tgctttttct tgccttatta ttgattagca agaaggaaat
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    46 caaattcata attattggtt taaccatgta atagtgcata ttaattgtga taaaaccttg
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    55 <212> TYPE: PRT
    56 <213> ORGANISM: Beta vulgaris
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    64 Ile Met Lys Gln Asn Ile Pro Glu Tyr Ser Leu Arg Phe Phe Ser Ile
    65
                                        25
    68 Ile Leu Glu Ile Ala Pro Ala Ala Lys Asn Met Phe Ser Phe Leu Arg
                                    40
    72 Asp Ser Glu Glu Val Pro Gln Asn Asn Pro Lys Leu Lys Ala His Ala
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76 Ile Lys Val Phe Lys Met Thr Cys Glu Ser Ala Ile Gln Leu Arg Glu

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77 65
                       70
                                           75
80 Lys Gly Glu Val Val Gly Glu Thr Thr Leu Lys Tyr Leu Gly Ala
84 Ile His Leu Lys Asn Gly Val Ile Asp Pro His Phe Glu Val Val Lys
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                                   105
88 Gln Ala Leu Leu Arg Thr Ile Glu Glu Ala Ser Gly Asp Lys Trp Ser
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92 Glu Glu Leu Lys Cys Ala Trp Ser Val Ala Tyr Asp His Leu Ala Ala
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96 Ala Ile Lys Ala Glu Met Lys Glu
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101 <211> LENGTH: 858
102 <212> TYPE: DNA
103 <213> ORGANISM: Arabidopsis thaliana
105 <400> SEQUENCE: 3
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108 ttcaaactgt gagagaaaaa gaaagagaga aagagatggg agagattggg tttacagaga
110 agcaagaagc tttggtgaag gaatcgtggg agatactgaa acaagacatc cccaaataca
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112 gccttcactt cttctcacag atactggaga tagcaccagc agcaaaaaggc ttgttctctt
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114 toctaagaga otcagatgaa gtoottoaca acaatootaa actoaaagot catgotgtta
116 aagtetteaa gatgacatgt gaaacageta tacagetgag ggaggaagga aaggtggtag
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118 tggctgacac aaccctccaa tatttaggct caattcatct caaaagcggc gttattgacc
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120 ctcacttcga ggtggtgaaa gaagctttgc taaggacatt gaaagagggg ttgggggaga
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122 aatacaatga agaagtggaa ggtgcttggt ctcaagctta tgatcacttg gctttagcca
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124 tcaagaccga gatgaaacaa gaagagtcat aaaaccctat tgatcatttg ggtatcgcat
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126 acatgaatet attecacata catgatacae atatacgtgt ttetgtgtgt gtactatgtt
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128 gctctctgac tttctacagt tcactatttt aattataaag aaggatcttg tgctatcatt
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130 agggagatac qtgatactgt agttcttctt gaaattgtta ttcgtgagaa atatcatggt
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132 ttgaaqtatt tattttcaca agatggatgt taacgtgggg atcattttac aatcattcta
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134 caaataattt tacttctc
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148 Ser Trp Glu Ile Leu Lys Gln Asp Ile Pro Lys Tyr Ser Leu His Phe
149
                20
152 Phe Ser Gln Ile Leu Glu Ile Ala Pro Ala Ala Lys Gly Leu Phe Ser
            35
                                40
156 Phe Leu Arg Asp Ser Asp Glu Val Pro His Asn Asn Pro Lys Leu Lys
160 Ala His Ala Val Lys Val Phe Lys Met Thr Cys Glu Thr Ala Ile Gln
161 65
164 Leu Arg Glu Glu Gly Lys Val Val Ala Asp Thr Thr Leu Gln Tyr
168 Leu Gly Ser Ile His Leu Lys Ser Gly Val Ile Asp Pro His Phe Glu
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172 Val Val Lys Glu Ala Leu Leu Arg Thr Leu Lys Glu Gly Leu Gly Glu
173
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                                120
176 Lys Tyr Asn Glu Glu Val Glu Gly Ala Trp Ser Gln Ala Tyr Asp His
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180 Leu Ala Leu Ala Ile Lys Thr Glu Met Lys Gln Glu Glu Ser
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                        150
181 145
184 <210> SEQ ID NO: 5
185 <211> LENGTH: 486
186 <212> TYPE: DNA
187 <213> ORGANISM: Brassica napus
189 <400> SEQUENCE: 5
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192 ctaaaqcaag atatccccaa atacagtett caettettet cacagatact ggagatagca
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194 ccaqcagcaa aqqacatqtt ctctttccta agagacacag atgaagtccc tcataacaat
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196 cctaaactca aagctcatgc tgttaaagtc ttcaagatga catgtgagac agcaatacag
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198 ctgagggaga aaggaaaggt agtggtggct gacacaaccc tccaatactt gggctctgtt
                                                                          300
200 catttcaaga geggtgttet tgateeteac tttgaggtgg tgaaagagge attggtgagg
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202 acactgaaag aagggttggg ggagaagtac aatgaagaag tggaaggagc ttggtccaag
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204 gcttatgatc acttggcttt agccattaag gccgagatga aacaagaaga ctcacaaaaa
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206 ccctaa
209 <210> SEQ ID NO: 6
210 <211> LENGTH: 161
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220 Ser Trp Glu Ile Leu Lys Gln Asp Ile Pro Lys Tyr Ser Leu His Phe
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224 Phe Ser Gln Ile Leu Glu Ile Ala Pro Ala Ala Lys Asp Met Phe Ser
225
            35
228 Phe Leu Arg Asp Thr Asp Glu Val Pro His Asn Asn Pro Lys Leu Lys
                             55
232 Ala His Ala Val Lys Val Phe Lys Met Thr Cys Glu Thr Ala Ile Gln
                                             75
236 Leu Arg Glu Lys Gly Lys Val Val Ala Asp Thr Thr Leu Gln Tyr
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                                         90
240 Leu Gly Ser Val His Phe Lys Ser Gly Val Leu Asp Pro His Phe Glu
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                                     105
                                                         110
244 Val Val Lys Glu Ala Leu Val Arg Thr Leu Lys Glu Gly Leu Gly Glu
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                                                     125
        . 115
248 Lys Tyr Asn Glu Glu Val Glu Gly Ala Trp Ser Lys Ala Tyr Asp His
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253 145
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256 Pro
260 <210> SEQ ID NO: 7
261 <211> LENGTH: 50
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Input Set : A:\CD054PCT.ST25.txt

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262 <212> TYPE: DNA 263 <213> ORGANISM: Artificial sequence 265 <220> FEATURE: 266 <223> OTHER INFORMATION: primer prm05458 268 <400> SEQUENCE: 7 50 269 ggggaccact ttgtacaaga aagctgggtc aaatgatcaa tagggtttta 272 <210> SEQ ID NO: 8 273 <211> LENGTH: 57 274 <212> TYPE: DNA 275 <213> ORGANISM: Artificial sequence 277 <220> FEATURE: 278 <223> OTHER INFORMATION: primer prm06122 280 <400> SEOUENCE: 8 281 ggggacaagt ttgtacaaaa aagcaggctt aaacagtgag agaaaaagaa agagaga 57 284 <210> SEQ ID NO: 9 285 <211> LENGTH: 54 286 <212> TYPE: DNA 287 <213> ORGANISM: Artificial sequence 289 <220> FEATURE: 290 <223> OTHER INFORMATION: primer prm05447 292 <400> SEQUENCE: 9 293 ggggacaagt ttgtacaaaa aagcaggctt aaacaatggc tctcgtggag gata 54 296 <210> SEQ ID NO: 10 297 <211> LENGTH: 48 298 <212> TYPE: DNA 299 <213> ORGANISM: Artificial sequence 301 <220> FEATURE: 302 <223> OTHER INFORMATION: primer prm05448 304 <400> SEQUENCE: 10 305 ggggaccact ttgtacaaga aagctgggtg atcatggagg tggagcag 48 308 <210> SEQ ID NO: 11 309 <211> LENGTH: 64 310 <212> TYPE: DNA 311 <213> ORGANISM: Artificial sequence 313 <220> FEATURE: 314 <223> OTHER INFORMATION: primer prm06021 316 <400> SEQUENCE: 11 317 ggggacaagt ttgtacaaaa aagcaggctt aaacaatgac ttttacagag aaagatgaag 60 319 cttt 64 322 <210> SEQ ID NO: 12 323 <211> LENGTH: 56 324 <212> TYPE: DNA 325 <213> ORGANISM: Artificial sequence 327 <220> FEATURE: 328 <223> OTHER INFORMATION: primer prm06022 330 <400> SEQUENCE: 12 331 ggggaccact ttgtacaaga aagctgggtc taagctacct attccttcat ctcagc

VERIFICATION SUMMARY

DATE: 10/14/2005 TIME: 13:27:42 PATENT APPLICATION: US/10/551,699

Input Set : A:\CD054PCT.ST25.txt

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L:9 M:270 C: Current Application Number differs, Replaced Current Application No L:9 M:271 C: Current Filing Date differs, Replaced Current Filing Date